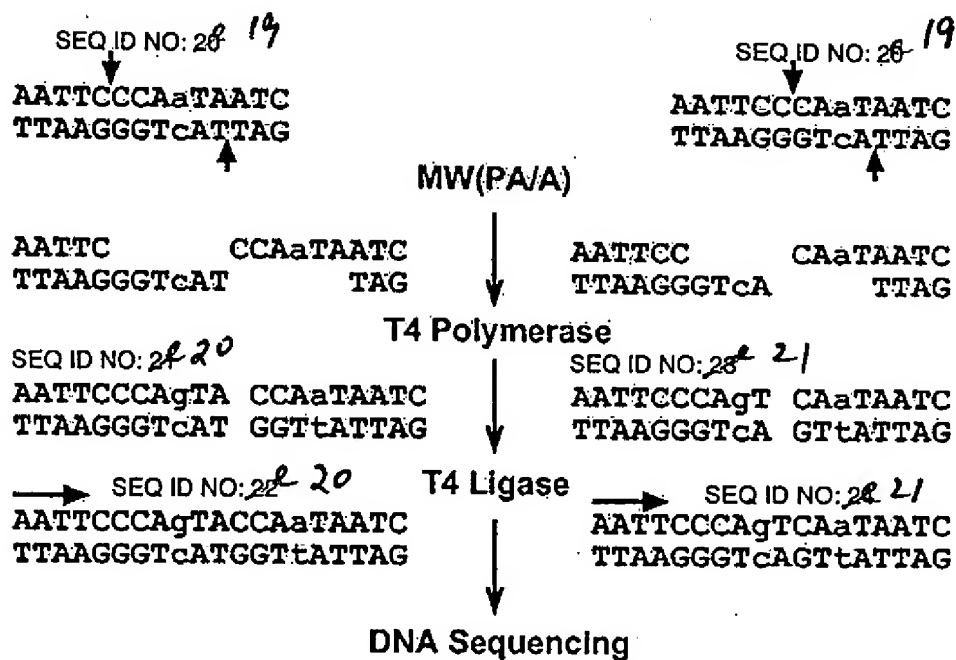


ANNOTATED DRAWING

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Figure 11 C



ANNOTATED DRAWING

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Figure 13

T7_endo1

>gi|431187:10257-10706 gene 3, endonuclease

ATGGCAGGTTACGGCGCTAAAGGAATCCGAAAGGTTGGAGCGTTTCGCTCTGGCC
TAGAGGACAAGGTTTTCAAAGCAGTTGGAATCAAAAGGTATTAAATTCGAGTATGAA
GAGTGGAAAGTGCCTTATGTAATTCCGGCGAGCAATCACACTTACACTCCAGACTT
CTTACTTCCAAACGGTATATTCGTTGAGACAAAGGGTCTGTGGGAAAGCGATGATA
GAAAGAAGCACTTATTAATTAGGGAGCAGCACCCCGAGCTAGACATCCGTATTGTC
TTCTCAAGCTCACGTACTAAGTTATACAAAGGTTCTCCAACGTCTTATGGAGAGTTC
TGCGAAAAGCATGGTATTAAGTTCGCTGATAAACTGATACCTGCTGAGTGGATAAA
GGAACCCAAGAAGGAGGTCCCCTTTGATAGATTAAAAAGGAAAGGAGGAAAGAAA
TAA (SEQ ID NO:1)

>gi|15581|emb|CAA24402.1| unnamed protein product [Enterobacteria
phage T7]

MAGYGAKGIRKVGAFRSGLEDKVSKQLESKGIKFYEYEWKVPYVIPASNHTYTPDFL
LPNGIFVETKGLWESDDRKKHLLIREQHPELDIRIVFSSSR TKLYKGSPTS YGEFCEK
HGIKFADKLIPAEWIKEPKKEVPFDR LKRKGKK (SEQ ID NO:15)

12

ANNOTATED DRAWING

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Figure 14-2

>gi|15517|emb|CAA24345.1| unnamed protein product

[Enterobacteria phage T7]

MAGYGAKGIRKVGAFRSGLEDKVSKQLESKGIKFEYEEWKVPYVIPASNHTYT
PDFLLPNGIFVETKGLWESDDRKKHLLIREQHPELDİRIVFSSSRTKLYKGSPTS
YGEFCEKHGKIFADKLIPAEWIKEPKKEVPFDRLKRKGGKK (SEQ ID NO:12)

=====

>gi|37956869|gb|AAP34135.1| gene 3 [Enterobacteria phage T7]

MAGYSAKGIRKVGAFRSGLEDKVSKQLESKGIKFEYEEWKVPYVIPASNHTYT
PDFLLPNGIFVETKGLWESDDRKKHLLIRKQHPELDİRIVFSSSRTKLYKGSPTS
YGEFCEKHGKIFADKLIPAEWIKEPKKEVPFDRLKRKGGKK (SEQ ID NO:24)
22

>gi|37956815|gb|AAP34082.1| gene 3 [Enterobacteria phage T7]

MAGYSAKGIRKVGAFRSGLEDKVSKQLESKGIKFEYEEWKVPYVIPASNHTYT
PDFLLPNGIFVETKGLWESDDRKKHLLIRKQHPELDİRIVFSSSRTKLYKGSPTS
YGEFCEKHGKIFADKLIPAEWIKEPKKEVPFDRLKRKGGKK (SEQ ID NO:24)
22

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>gi|37956764|gb|AAP34032.1| gene 3 [Enterobacteria phage T7]

MVGYGKVGIRKVGAFRSGLEDKVSKQLESKGIKFEYEEWKVPYVIPASNHTYT
PDFLLPNGIFVETKGLWESDDRKKHLLIREQHPELDİRIVFSSSRTKLYKGSPTS
YGEFCEKHGKIFADKLIPAEWIKEPKKEVSFDRLKRKGGKK (SEQ ID NO:25)
23

ANNOTATED DRAWING

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Figure 14-3

>gi|37956712|gb|AAP33981.1| gene 3 [Enterobacteria phage T7]
MVGYGVKGIRKVGAFRSGLEDKVSKQLESKGIKFEYEEWKVPYVIPASNHTYT
PDFLLPNGIFVETKGLWESDDRKKHLLIREQHPELDIRIVFSSSRTKLYKGSPTS
YGEFCEKHGKIFADKLIPAEWIKPKKEVSFDRLKRKGGKK (SEQ ID NO:28)

23

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>gi|30387466|ref|NP_848275.1| endonuclease [Yersinia pestis phage
phiA1122]
MAGTYAARGIRKVGTFRSGLEDKVSKQLEGKGKIKFDYELWKIPYVVPASNHVY
TPDFLLPNGIFIETKGLWESDDRKKHLLIREQFPELDRLVFSSSRTKLYKGSPT
SYGEWCEKHGILFADKLIPVEWLKEPKKEVPFDRLKQAKGGKK
(SEQ ID NO:13)

>gi|30314103|gb|AAP20511.1| endonuclease [Yersinia pestis phage
phiA1122]
MAGTYAARGIRKVGTFRSGLEDKVSKQLEGKGKIKFDYELWKIPYVVPASNHVY
TPDFLLPNGIFIETKGLWESDDRKKHLLIREQFPELDRLVFSSSRTKLYKGSPT
SYGEWCEKHGILFADKLIPVEWLKEPKKEVPFDRLKQAKGGKK
(SEQ ID NO:13)

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